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5. (Amended) A support as claimed in claim 1, wherein each oligonucleotide is non-complementary with genomic DNA and non complementary with each other.

9. (Amended) A method as claimed in claim 7, wherein each primer is used in conjunction with a second companion primer to amplify the target region of interest.

12. (Amended) A method as claimed in claim 10, wherein there are between 5 and 80 distinct targeting oligonucleotides per target gene.

13. (Amended) A method as claimed in claim 10, wherein step (iii) is carried out as separate reactions between the target cDNAs and separate pools of targeting polynucleotides, the cDNA target binding sequences of the polynucleotides in each pool possessing approximately the same T_m as the others in the pool; the hybridisation reactions from each pool of targeting polynucleotides are then pooled together.--